

ABSTRACT OF THE DISCLOSURE

Method for determining a nucleotide sequence of
an analytical oligo nucleic acid for use in analysis of
a nucleic acid, comprising the steps of listing all
unit nucleotide sequences present on a target nucleic
acid having a predetermined length which is shorter
than that of the analytical oligo nucleic acid to
be designed, and extracting a nucleotide sequence
containing a sequence occurring at a low frequency
on the target nucleic acid from the candidate sequences
of the analytical oligo nucleic acid on the basis of
a frequency of each of the listed individual unit
sequences.

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